ELSEVIER

Gene 174 (1996) 323–325

GENE
AN INTERNATIONAL JOURNAL ON GENES AND GENOMES

Author Index

Abe, M., Tsuda, M., Kimoto, M., Inouye, S., Nakazawa, A. and Nakazawa, T.
A genetic analysis system of *Burkholderia cepacia*: construction of mobilizable transposons and a cloning vector (174) 191
Ahmad, D., see Sylvestre, M. (174) 195
Anderson, O.D., Kuhl, J.C. and Tam, A.
Construction and expression of a synthetic wheat storage protein gene (174) 51
Andreozzi, L., see Saccone, S. (174) 85
Apostolou, S., see Scott, I.C. (174) 135
Arber, W., see Szeverényi, I. (174) 103
Aszodi, A., see Baranyi, M. (174) 27
Augé-Gouillou, C., see Bigot, Y. (174) 265
Avalos, J., Corrochano, L.M. and Brenner, S.
Genomic organization of the fungus *Phycomyces* (174) 43

Baranyi, M., Aszodi, A., Devinoy, E., Fontaine, M.-L., Houdebine, L.-M. and Bösze, Z.
Structure of the rabbit κ-casein encoding gene: expression of the cloned gene in the mammary gland of transgenic mice (174) 27
Barriault, D., see Sylvestre, M. (174) 195
Beckmann, J.D., see Henry, T. (174) 221
Beggs, M.L., Cave, M.D. and Eisenach, K.D.
Isolation and sequence of a *Mycobacterium tuberculosis* sigma factor (174) 285
Bègue, A., see Grévin, D. (174) 185
Beh, K.J., see Callaghan, M.J. (174) 273
Bentle, K.A., see Warren, W.C. (174) 235
Bergeron, J., see Sylvestre, M. (174) 195
Bernardi, G., see Saccone, S. (174) 85
Bernardi, G., see Zoubak, S. (174) 95
Betz, A., see Mouchel, N. (174) 59
Bhat-Nakshatri, P., see Nakshatri, H. (174) 79
Bigot, Y., Augé-Gouillou, C. and Periquet, G.
Computer analyses reveal a *hobo*-like element in the nematode *Caenorhabditis elegans*, which presents a conserved transposase domain common with the Tc1-*Mariner* transposon family (174) 265
Billaut-Mulot, O., Fernandez-Gomez, R., Loyens, M. and Ouaissi, A.
Trypanosoma cruzi elongation factor 1-α: nuclear localization in parasites undergoing apoptosis (174) 19
Blackshear, P.J., see Thompson, M.J. (174) 225
Boavida, M.G., see Marshall, B. (174) 175
Bodescot, M. and Brison, O.
Characterization of new human c-*myc* mRNA species produced by alternative splicing (174) 115
Bogosian, G., see Warren, W.C. (174) 235
Bosworth, B.T., see Lee, E.-k. (174) 311
Bösze, Z., see Baranyi, M. (174) 27
Bouillet, P., see Nakshatri, H. (174) 79
Bram, R.J.
Isolation of a cDNA encoding the chicken homologue of the human calcium-modulating cyclophilin ligand (174) 307
Bréant, C., see Eichmann, A. (174) 3
Brenner, S., see Avalos, J. (174) 43
Brison, O., see Bodescot, M. (174) 115
Brown, J.C.S., see Ward, L.J.H. (174) 259

Cacciò, S., see Saccone, S. (174) 85
Callaghan, M.J. and Beh, K.J.
A tandemly repetitive DNA sequence is present at diverse locations in the genome of *Ostertagia circumcincta* (174) 273
Callen, D.F., see Scott, I.C. (174) 135
Cave, M.D., see Beggs, M.L. (174) 285
Chambon, P., see Nakshatri, H. (174) 79
Chang, W.-C., see Hsu, L.C. (174) 319
Chatterjee, B., see Zarbalis, K. (174) 181
Chen, D., Guo, J., Miki, T., Tachibana, M. and Gahl, W.A.
Molecular cloning of two novel *rab* genes from human melanocytes (174) 129
Chen, H.-Y., see Wu, C.-H. (174) 251
Chin, H., see Ratty, A.K. (174) 71
Clay, O., see Zoubak, S. (174) 95
Collart, F.R., Osipiuk, J., Trent, J., Olsen, G.J. and Huberman, E.
Cloning, characterization and sequence comparison of the gene coding for IMP dehydrogenase from *Pyrococcus furiosus* (174) 209
Collart, F.R., Osipiuk, J., Trent, J., Olsen, G.J. and Huberman, E.
Cloning and characterization of the gene encoding IMP dehydrogenase from *Arabidopsis thaliana* (174) 217
Corrochano, L.M., see Avalos, J. (174) 43

Davey, G.P., see Ward, L.J.H. (174) 259
De Macario, E.C., see Jovell, R.J. (174) 281
Dehem, M., see Grévin, D. (174) 185
Denhez, F., see Grévin, D. (174) 185
Devinoy, E., see Baranyi, M. (174) 27
Dietz, A.B., see Lee, E.-k. (174) 311
Dujon, B., see Richard, G.-F. (174) 165
Dumas, B., see Lacour, T. (174) 289

Edlund, A., Johansson, T., Leidvik, B. and Hansson, L.
Structure of the human κ-casein gene (174) 65
Eichmann, A., Marcelle, C., Bréant, C. and Le Douarin, N.M.
Molecular cloning of Quek 1 and 2, two quail vascular endothelial growth factor (VEGF) receptor-like molecules (174) 3
Eisenach, K.D., see Beggs, M.L. (174) 285
Entwistle, J., see Wang, C. (174) 299
Erttmann, K.D. and Gallin, M.Y.
Onchocerca volvulus: identification of cDNAs encoding a putative phosphatidyl-ethanolamine-binding protein and a putative partially processed mRNA precursor (174) 203

Farris, A.D., Gross, J.K., Hanas, J.S. and Harley, J.B.
Genes for murine Y1 and Y3 Ro RNAs have Class 3 RNA polymerase III promoter structures and are unlinked on mouse chromosome 6 (174) 35
Feix, G., see Goddemeier, M.L. (174) 111
Feoktistova, A., see Ohi, R. (174) 315
Fernandez-Gomez, R., see Billaut-Mulot, O. (174) 19
Fontaine, M.-L., see Baranyi, M. (174) 27
Forterre, P., see Guipaud, O. (174) 121
Funkenstein, B. and Jakowlew, S.B.
Molecular cloning of fish alcohol dehydrogenase cDNA (174) 159

Gahl, W.A., see Chen, D. (174) 129
Gainer, H., see Ratty, A.K. (174) 71
Gallin, M.Y., see Erttmann, K.D. (174) 203
Geryk, J., see Machoň, O. (174) 9
Goddemeier, M.L. and Feix, G.
Genomic structure of the maize TATA-box binding protein 1 (TBP-1): conserved exon/intron structure in eukaryotic *TBP* genes (174) 111
Gould, K.L., see Ohi, R. (174) 315
Graw, J., see Zarbalis, K. (174) 181
Grévin, D., Ung, S., Denhez, F., Dehem, M., Quatannens, B., Bègue, A., Stéhelin, D. and Martin, P.
Structure and organization of the mouse *elk1* gene (174) 185
Gross, J.K., see Farris, A.D. (174) 35
Guillemette, I., see Sylvestre, M. (174) 195
Guipaud, O., Labedan, B. and Forterre, P.
A *gyrB*-like gene from the hyperthermophilic bacterion *Thermotoga maritima* (174) 121
Guo, J., see Chen, D. (174) 129

Hájková, P., see Machoň, O. (174) 9
Halila, R., see Scott, I.C. (174) 135
Hanas, J.S., see Farris, A.D. (174) 35
Hansson, L., see Edlund, A. (174) 65
Harley, J.B., see Farris, A.D. (174) 35
Heinaru, A., see Kõiv, V. (174) 293
Hejnar, J., see Machoň, O. (174) 9
Henry, T., Kliewer, B., Palmatier, R., Ulphani, J.S. and Beckmann, J.D.
Isolation and characterization of a bovine gene encoding phenol sulfotransferase (174) 221
Hodel, A., see Szeverényi, I. (174) 103
Hou, G., see Wang, C. (174) 299
Houdebine, L.-M., see Baranyi, M. (174) 27
Hsu, L.C. and Chang, W.-C.
Sequencing and expression of the human *ALDH8* encoding a new member of the aldehyde dehydrogenase family (174) 319
Huberman, E., see Collart, F.R. (174) 209
Huberman, E., see Collart, F.R. (174) 217
Hurtubise, Y., see Sylvestre, M. (174) 195

Inouye, S., see Abe, M. (174) 191
Isidro, G., see Marshall, B. (174) 175

Jakowlew, S.B., see Funkenstein, B. (174) 159
Jenkins, J.M., see Scott, I.C. (174) 135
Jeong, S.-W., see Ratty, A.K. (174) 71
Johansson, T., see Edlund, A. (174) 65
Jovell, R.J., Macario, A.J.L. and Macario, E.C.d.
ABC transporters in Archaea: two genes encoding homologs of the nucleotide-binding components in the methanogen *Methanosarcina mazei* S-6 (174) 281
Juteau, J.M., see Sylvestre, M. (174) 195

Kadler, K.E., see Scott, I.C. (174) 135
Kang, D., Romans, P. and Lee, J.-Y.
Analysis of a lysozyme gene from the malaria vector mosquito, *Anopheles gambiae* (174) 239
Kang, D., Lundström, A. and Steiner, H.
Trichoplusia ni attacin A, a differentially displayed insect gene coding for an antibacterial protein (174) 245
Kawamoto, S., Matsumoto, Y., Mizuno, K., Okubo, K. and Matsubara, K.
Expression profiles of active genes in human and mouse livers (174) 151
Kehrli, M.E. Jr., see Lee, E.-k. (174) 311
Kimoto, M., see Abe, M. (174) 191
Kliewer, B., see Henry, T. (174) 221
Kõiv, V., Marits, R. and Heinaru, A.
Sequence analysis of the 2,4-dichlorophenol hydroxylase gene *tfdB* and 3,5-dichlorocatechol 1,2-dioxygenase gene *tfdC* of 2,4-dichlorophenoxyacetic acid degrading plasmid pEST4011 (174) 293
Kuhl, J.C., see Anderson, O.D. (174) 51
Kusuda, J., see Saccone, S. (174) 85

Labedan, B., see Guipaud, O. (174) 121
Lacour, T. and Dumas, B.
A gene encoding a yeast equivalent of mammalian NADPH-adrenodoxin oxidoreductases (174) 289
Lai, Wi.S., see Thompson, M.J. (174) 225
Le Douarin, N.M., see Eichmann, A. (174) 3
Le Pennec, J.P., see Mouchel, N. (174) 59
Lee, E.-k., Kehrli, M.E. Jr., Dietz, A.B., Bosworth, B.T. and Reinhardt, T.A.
Cloning and sequencing of a cDNA encoding bovine intercellular adhesion molecule 3 (ICAM-3)[1] (174) 311
Lee, J.-Y., see Kang, D. (174) 239
Leidvik, B., see Edlund, A. (174) 65
Li, Q., see Wang, C. (174) 299
Löster, J., see Zarbalis, K. (174) 181
Loyens, M., see Billaut-Mulot, O. (174) 19
Lundström, A., see Kang, D. (174) 245

Macario, A.J.L., see Jovell, R.J. (174) 281
Machoň, O., Hejnar, J., Hájková, P., Geryk, J. and Svoboda, J.
The LTR, v-*src*, LTR provirus in H-19 hamster tumor cell line is integrated adjacent to the negative regulatory region (174) 9
Marcelle, C., see Eichmann, A. (174) 3
Marits, R., see Kõiv, V. (174) 293
Marshall, B., Isidro, G. and Boavida, M.G.
Insertion of a short Alu sequence into the hMSH2 gene following a double cross over next to sequences with chi homology (174) 175
Martin, P., see Grévin, D. (174) 185
Matsubara, K., see Kawamoto, S. (174) 151
Matsumoto, Y., see Kawamoto, S. (174) 151
Mehan, S., see Scott, I.C. (174) 135
Miki, T., see Chen, D. (174) 129
Mizuno, K., see Kawamoto, S. (174) 151
Mouchel, N., Trichet, V., Betz, A., Pennec, J.P. and Wolff, J.
Characterization of vitellogenin from rainbow trout (*Oncorhynchus mykiss*) (174) 59
Mullany, P., Pallen, M., Wilks, M., Stephen, J.R. and Tabaqchali, S.
A Group II intron in a conjugative transposon from the Gram-positive bacterium, *Clostridium difficile* (174) 145
Murphy, D., see Ratty, A.K. (174) 71

Nagle, J.W., see Ratty, A.K. (174) 71
Nakazawa, A., see Abe, M. (174) 191
Nakazawa, T., see Abe, M. (174) 191

Nakshatri, H., Bouillet, P., Bhat-Nakshatri, P. and Chambon, P.
Isolation of retinoic acid-repressed genes from P19 embryonal carcinoma cells (174) 79

O'Neil, J.P., see Warren, W.C. (174) 235
Ohi, R., Feoktistova, A. and Gould, K.L.
Construction of vectors and a genomic library for use with *his3*-deficient strains of *Schizosaccharomyces pombe* (174) 315
Okubo, K., see Kawamoto, S. (174) 151
Olasz, F., see Szeverényi, I. (174) 103
Olsen, G.J., see Collart, F.R. (174) 209
Olsen, G.J., see Collart, F.R. (174) 217
Osipiuk, J., see Collart, F.R. (174) 209
Osipiuk, J., see Collart, F.R. (174) 217
Ouaissi, A., see Billaut-Mulot, O. (174) 19

Pallen, M., see Mullany, P. (174) 145
Palmatier, R., see Henry, T. (174) 221
Peltonen, L., see Scott, I.C. (174) 135
Periquet, G., see Bigot, Y. (174) 265
Prockop, D.J., see Scott, I.C. (174) 135

Quatannens, B., see Grévin, D. (174) 185

Ratty, A.K., Jeong, S.-W., Nagle, J.W., Chin, H., Gainer, H., Murphy, D. and Venkatesh, B.
A systematic survey of the intergenic region between the murine oxytocin- and vasopressin-encoding genes (174) 71
Reinhardt, T.A., see Lee, E.-k. (174) 311
Richard, G.-F. and Dujon, B.
Distribution and variability of trinucleotide repeats in the genome of the yeast *Saccharomyces cerevisiae* (174) 165
Romans, P., see Kang, D. (174) 239

Saccone, S., Cacciò, S., Kusuda, J., Andreozzi, L. and Bernardi, G.
Identification of the gene-richest bands in human chromosomes (174) 85
Schlittler, M.R., see Warren, W.C. (174) 235
Schwane, A.C., see Warren, W.C. (174) 235
Scott, I.C., Halila, R., Jenkins, J.M., Mehan, S., Apostolou, S., Winqvist, R., Callen, D.F., Prockop, D.J., Peltonen, L. and Kadler, K.E.
Molecular cloning, expression and chromosomal localization of a human gene encoding a 33 kDa putative metallopeptidase (PRSM1) (174) 135
Shareck, F., see Sylvestre, M. (174) 195
Shiuan, D., see Wu, C.-H. (174) 251
Sirois, M., see Sylvestre, M. (174) 195
Stéhelin, D., see Grévin, D. (174) 185
Steiner, H., see Kang, D. (174) 245
Stephen, J.R., see Mullany, P. (174) 145
Svoboda, J., see Machoň, O. (174) 9
Sylvestre, M., Sirois, M., Hurtubise, Y., Bergeron, J., Ahmad, D., Shareck, F., Barriault, D., Guillemette, I. and Juteau, J.M.
Sequencing of *Comamonas testosteroni* strain B-356-biphenyl/chlorobiphenyl dioxygenase genes: evolutionary relationships among Gram-negative bacterial biphenyl dioxygenases (174) 195
Szeverényi, I., Hodel, A., Arber, W. and Olasz, F.
Vector for IS element entrapment and functional characterization based on turning on expression of distal promoterless genes (174) 103

Tabaqchali, S., see Mullany, P. (174) 145
Tachibana, M., see Chen, D. (174) 129
Tam, A., see Anderson, O.D. (174) 51
Taylor, G.A., see Thompson, M.J. (174) 225
Thompson, M.J., Lai, Wi.S., Taylor, G.A. and Blackshear, P.J.
Cloning and characterization of two yeast genes encoding members of the CCCH class of zinc finger proteins: zinc finger-mediated impairment of cell growth (174) 225
Trent, J., see Collart, F.R. (174) 209
Trent, J., see Collart, F.R. (174) 217
Trichet, V., see Mouchel, N. (174) 59
Tsuda, M., see Abe, M. (174) 191
Turley, E.A., see Wang, C. (174) 299

Ulphani, J.S., see Henry, T. (174) 221
Ung, S., see Grévin, D. (174) 185

Venkatesh, B., see Ratty, A.K. (174) 71

Wang, C., Entwistle, J., Hou, G., Li, Q. and Turley, E.A.
The characterization of a human *RHAMM* cDNA: conservation of the hyaluronan-binding domains (174) 299
Ward, L.J.H., Brown, J.C.S. and Davey, G.P.
Identification and sequence analysis of IS*1297*, an IS*S1*-like insertion sequence in a *Leuconostoc* strain (174) 259
Warren, W.C., Bentle, K.A., Schlittler, M.R., Schwane, A.C., O'Neil, J.P. and Bogosian, G.
Increased production of peptide deformylase eliminates retention of formylmethionine in bovine somatotropin overproduced in *Escherichia coli* (174) 235
Werner, T., see Zarbalis, K. (174) 181
Wilks, M., see Mullany, P. (174) 145
Winqvist, R., see Scott, I.C. (174) 135
Wolff, J., see Mouchel, N. (174) 59
Wu, C.-H., Chen, H.-Y. and Shiuan, D.
Isolation and characterization of the *Erwinia herbicola bio* operon and the sequences of the *bioA* and *bioB* genes (174) 251

Zarbalis, K., Chatterjee, B., Löster, J., Werner, T. and Graw, J.
Sequence analysis of the βB2-crystallin cDNA of hamster containing a domain conserved among vertebrates (174) 181
Zoubak, S., Clay, O. and Bernardi, G.
The gene distribution of the human genome (174) 95